

coupledCahnHilliardAllenCahn tutorial

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1 Parameters

This section describes the model parameters that the user can access through the `parameters.h` file in each application folder. These can be divided between model-specific parameters and those generic to all models, and there are three categories of model-specific parameters: constant coefficients and tensors, bulk free energy functions, and residuals. In the context of their model, constant coefficients are self-explanatory. Bulk free energy functions and their derivatives are incorporated as functions of the phase field variables, usually instantiated in the code as `n` or `c`. All of the examples and defaults use polynomial free energies, and their derivatives are found symbolically. Non-polynomial functions could be implemented, but are not supported at this time. Residuals are used directly in evolution. Changing them is not recommended unless the user is comfortable with deal.ii and the PRISMS framework.

1.1 Generic Parameters

- **problemDIM**
Dimension of the problem (e.g. 1D, 2D, 3D)
- **spanX**
Length of system in x-direction
- **spanY**
Length of system in y-direction. Not used if `problemDIM < 2`
- **spanZ**
Length of system in z-direction. Not used if `problemDIM < 3`
- **refineFactor**
Defines the refinement of the mesh. There are $2^{\text{refineFactor}}$ elements in each direction in this implementation, and $\left(2^{\text{refineFactor}}\right)^{\text{problemDIM}}$ elements in total.
- **finiteElementDegree**
The order of interpolation of the finite element space. In this case, the order of the Lagrange elements to be used.

- **dt**
The simulation timestep.
- **numIncrements**
The number of simulation iterations. Final time is then `dt*numIncrements`.
- **writeOutput**
Whether we are writing any output. Takes a boolean argument, e.g. `true`.
- **skipOutputSteps**
Output will be written every `skipOutputSteps` iterations. If `writeOutput true`, the initial conditions will always be written.

1.2 coupledCahnHilliardAllenCahn Parameters

This application implements the following set of equations for a conserved order parameter, c , and one or more non-conserved (structural) order parameters η :

$$\frac{\partial \eta}{\partial t} = -M_\eta \left(\frac{\partial f(\eta, c)}{\partial \eta} - K_{ij} \nabla_i \nabla_j \eta \right) \quad (1)$$

$$\frac{\partial c}{\partial t} = \nabla \cdot M_c \nabla \left(\frac{\partial f(\eta, c)}{\partial c} \right) \quad (2)$$

Since there is no gradient energy term in c , this might also be called a coupled Allen-Cahn/diffusion system. From a finite element perspective, Eq. 5 is more tractable as

$$\frac{\partial c}{\partial t} = \nabla \cdot M_c \left(\frac{\partial^2 f(\eta, c)}{\partial c^2} \nabla c + \frac{\partial^2 f(\eta, c)}{\partial \eta \partial c} \nabla \eta \right) \quad (3)$$

- **numStructuralOrderParameters**
Multiple structural/non-conserved order parameters can be used in this model (implemented?????)
- **Mc**
 M_c , mobility of the conserved order parameter c .
- **MnVals**
 M_η , mobility of the non-conserved order parameter η . This is an array with `numStructuralOrderParameters` elements, each corresponding to an order parameter.
- **KnVals**
 K_η , gradient energy coefficients K of the non-conserved order parameter η . Again, an array with `numStructuralOrderParameters` elements, each corresponding to an order parameter.

The free energy $f(\eta, c)$ is composed of free energies $f_\alpha(c)$ and $f_\beta(c)$ for the phases corresponding to $\eta = 1$ and $\eta = 0$, and an interpolating function $h(\eta)$. The default case uses free energies for a Mg-Gd system:

$$f_\alpha(c) = -1.6704 - 4.776c + 5.1622c^2 - 2.7375c^3 + 1.3687c^4 \quad (4)$$

$$f_\beta(c) = -1.5924 - 5.9746c + 5.0c^2 \quad (5)$$

These free energies should be in the form of a polynomial for this application. An interpolation function for a single order parameter should satisfy $h(0) = 1$, $h'(0) = 0$, $h(1) = 1$, and $h'(1) = 0$, for example:

$$h(\eta) = \eta^3(10 - 15\eta + 6\eta^2) \quad (6)$$

The full bulk free energy (for n structural order parameters) is then:

$$f(\eta, c) = f_\alpha(c) \sum_{i=1}^n (1 - h(\eta_i)) + f_\beta(c) \sum_{i=1}^n h(\eta_i) \quad (7)$$

Equations (7-9) and their derivatives are included in `parameters.h` as follows:

$$\mathbf{faV} = f_\alpha(c)$$

$$\mathbf{facV} = f'_\alpha(c)$$

$$\mathbf{faccV} = f''_\alpha(c)$$

$$\mathbf{fbV} = f_\beta(c)$$

$$\mathbf{fbcV} = f'_\beta(c)$$

$$\mathbf{fbccV} = f''_\beta(c)$$

$$\mathbf{hV} = h(\eta)$$

$$\mathbf{hnV} = h'(\eta)$$

These free energy variables are incorporated into the residuals in the weak form of equations 4 and 6.

$$\mathbf{rcxV} = \left((1 - h(\eta))f''_\alpha(c) + h(\eta)f''_\beta(c) \right) \nabla c + h'(\eta)(f'_\beta(c) + f'_\alpha(c))\nabla \eta$$

$$\mathbf{rnV} = (f_\beta(c) - f_\alpha(c))h'(\eta)$$

$$\mathbf{rnxV} = \nabla \eta$$

2 Initial Conditions

Initial conditions for the phase field parameters are generated in the functions `InitialConditionN` or `InitialConditionC` located in the `main.cc` file in the application folder. Two types of initial conditions have been implemented so far: a field of random noise around some mean value, and analytical functions of position. Random noise is the default for at least one variable in all of the phase field codes. Here, as an example, c is initialized to random values in the range 0.019-0.021:

```
template <int dim>
double InitialConditionC<dim>::value (const Point<dim> &p, const unsigned int
    /* component */ const
{
    //set result equal to the concentration initial condition
    return 0.02 + 1.0e-3*(2*(0.5 - (double)(std::rand() % 100 )/100.0));
}
```

Analytical functions of position can be generated from the members of the class `p`, e.g. `p.distance` or the x and y coordinates, `p[0]` and `p[1]`. In the default cases, the hyperbolic tangent of the signed distance from a circle or sphere is used. In the following example, η will be initialized into a sphere with radius= $\text{spanX}/8$, where $\eta = 0.5$ at the boundary, $\eta \sim 1$ in the interior, and $\eta \sim 0$ outside.

```
template <int dim>
double InitialConditionN<dim>::value (const Point<dim> &p, const unsigned int
    /* component */ const
{
    //set result equal to the structural order paramter initial condition
    double dx=spanX/std::pow(2.0,refineFactor);
    double r=0.0;
    r=p.distance(Point<dim>(spanX/2.0,spanY/2.0,spanZ/2.0));
    return 0.5*(1.0-std::tanh((r-spanX/8.0)/(3*dx)));
}
```

This example requires that `problemDIM=3`. In the actual applications (e.g. `coupledCahnHilliardAllenCahn`), a conditional tests `problemDIM`, and there are analogous cases for each dimension.

3 Boundary Conditions

Boundary conditions are no-flux for the phase field evolution equations. Implementation of alternate boundary conditions is an objective for future releases.